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$$E = R^{-6}/(R^{-6} + R_0^{-6})$$

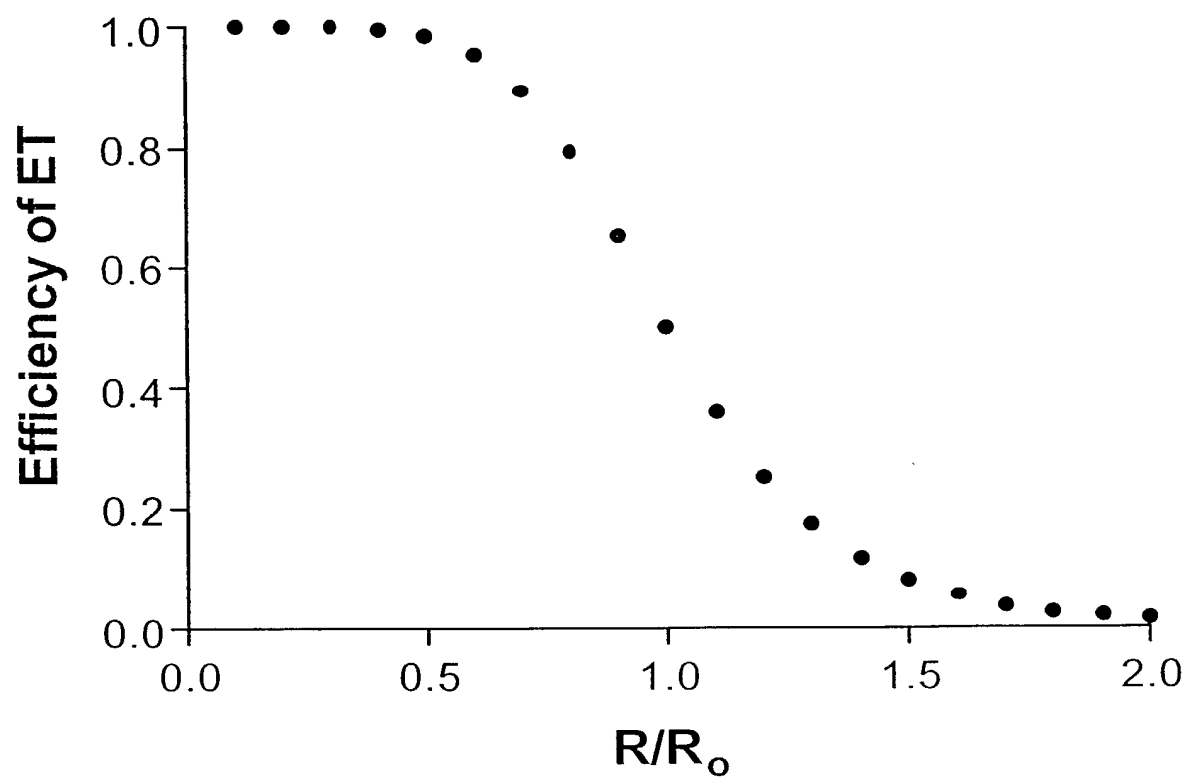


FIG. 1

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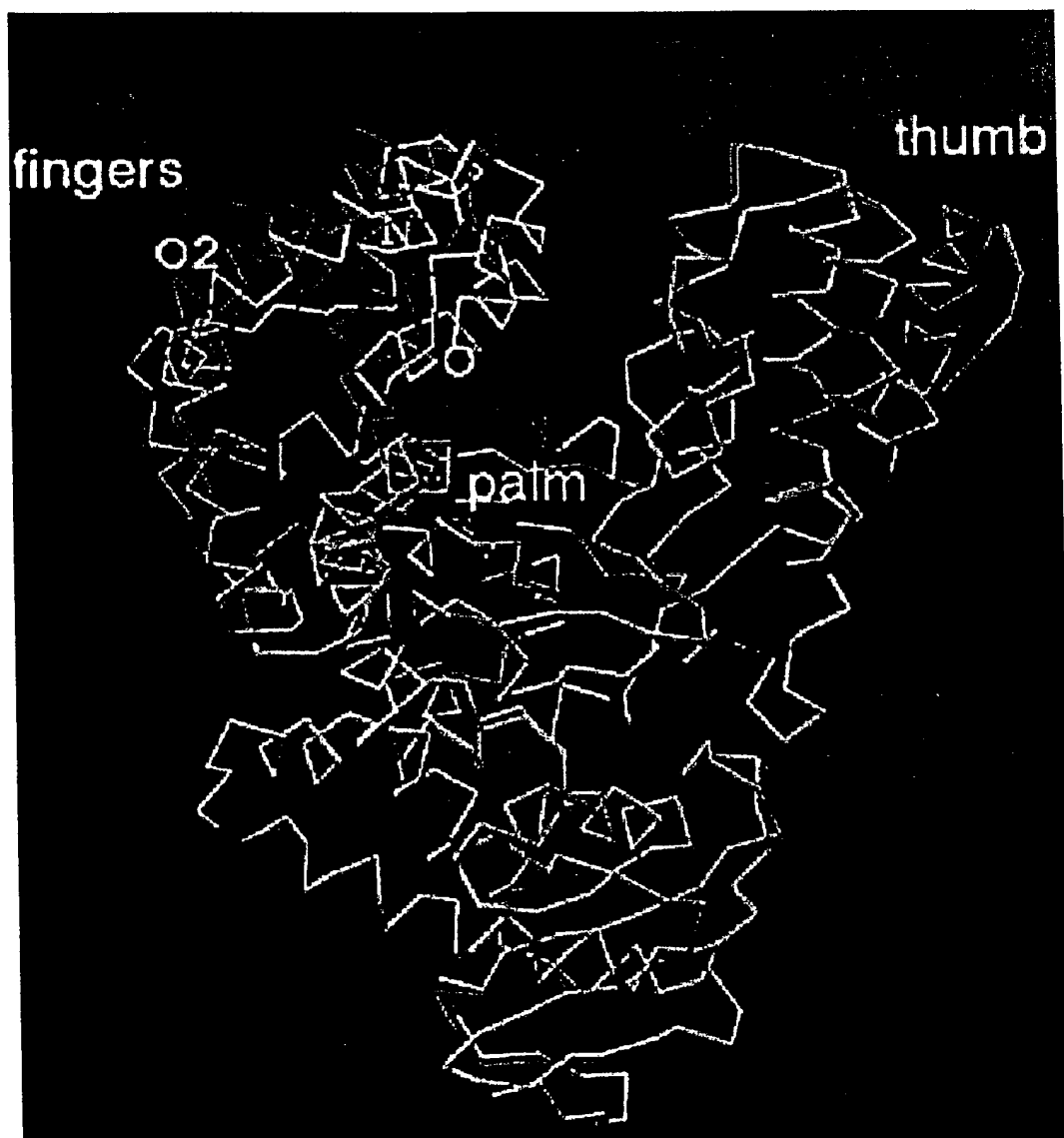


FIG. 2

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FIG. 3A

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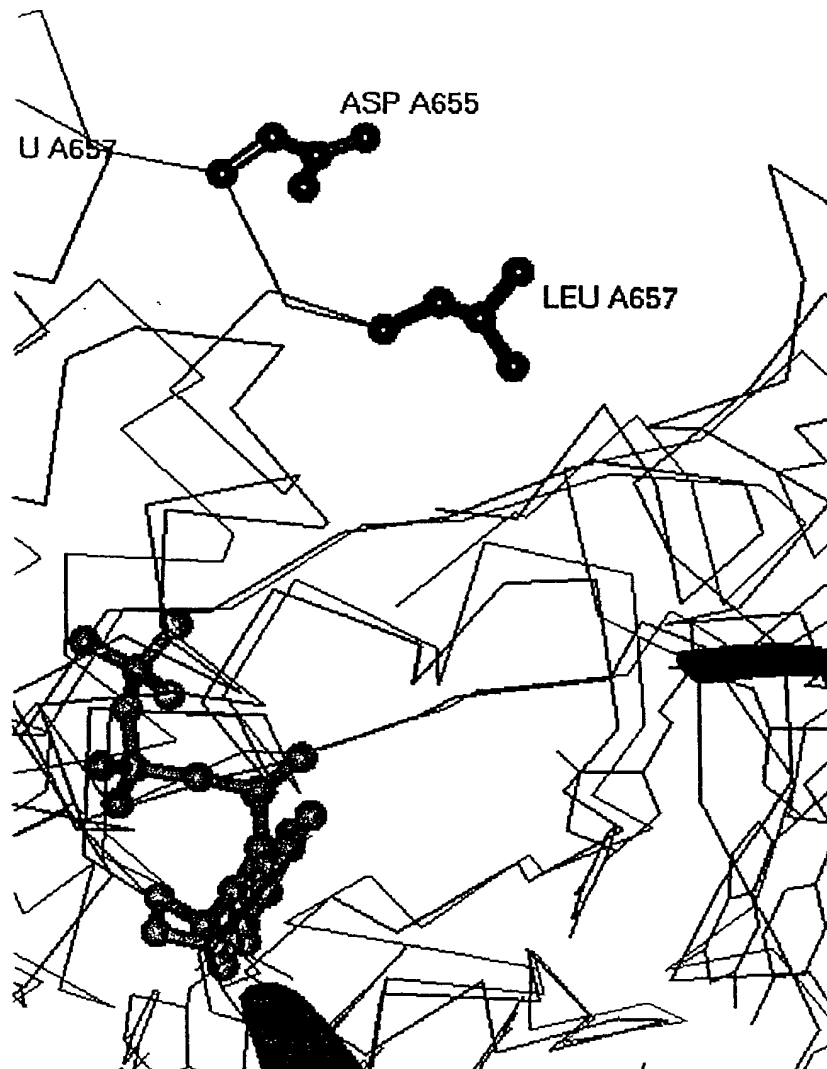


FIG. 3B

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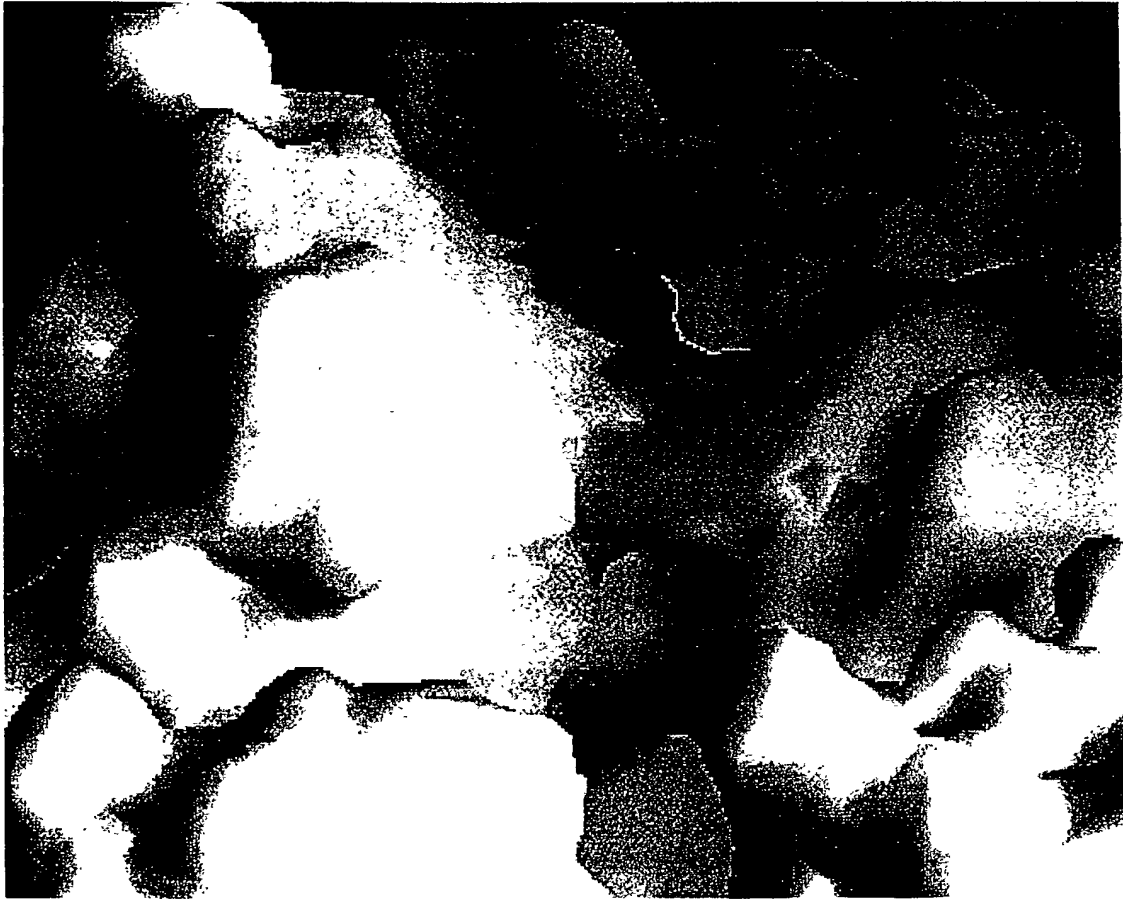


FIG. 3C

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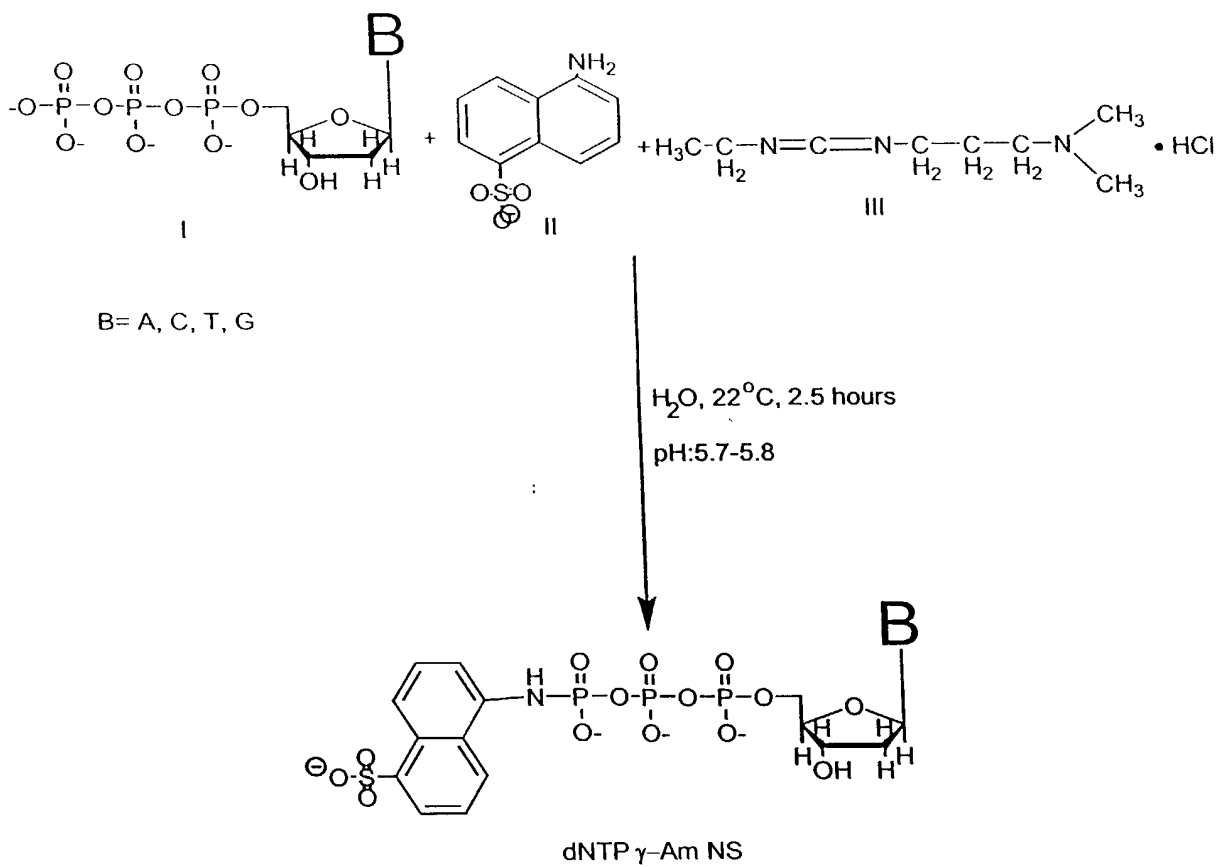


FIG. 4

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| | |
|-------------------|-----------------------------------|
| Primer Strand: | |
| TOP 5' | GGT ACT AAG CGG CCG CAT G 3' |
| Template Strands: | |
| BOT-T 3' | CCA TGA TTC GCC GGC GTA CTC 5' |
| BOT-C 3' | CCA TGA TTC GCC GGC GTA CCC 5' |
| BOT-G 3' | CCA TGA TTC GCC GGC GTA CGC 5' |
| BOT-A 3' | CCA TGA TTC GCC GGC GTA CAC 5' |
| BOT-3T 3' | CCA TGA TTC GCC GGC GTA CTT TC 5' |
| BOT-Sau 3' | CCA TGA TTC GCC GGC GTA CCT AG 5' |

Incorporate: GATC AG AAAG
(5' to 3')

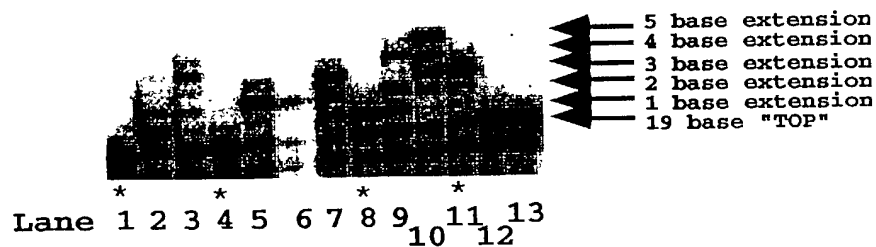


FIG. 5

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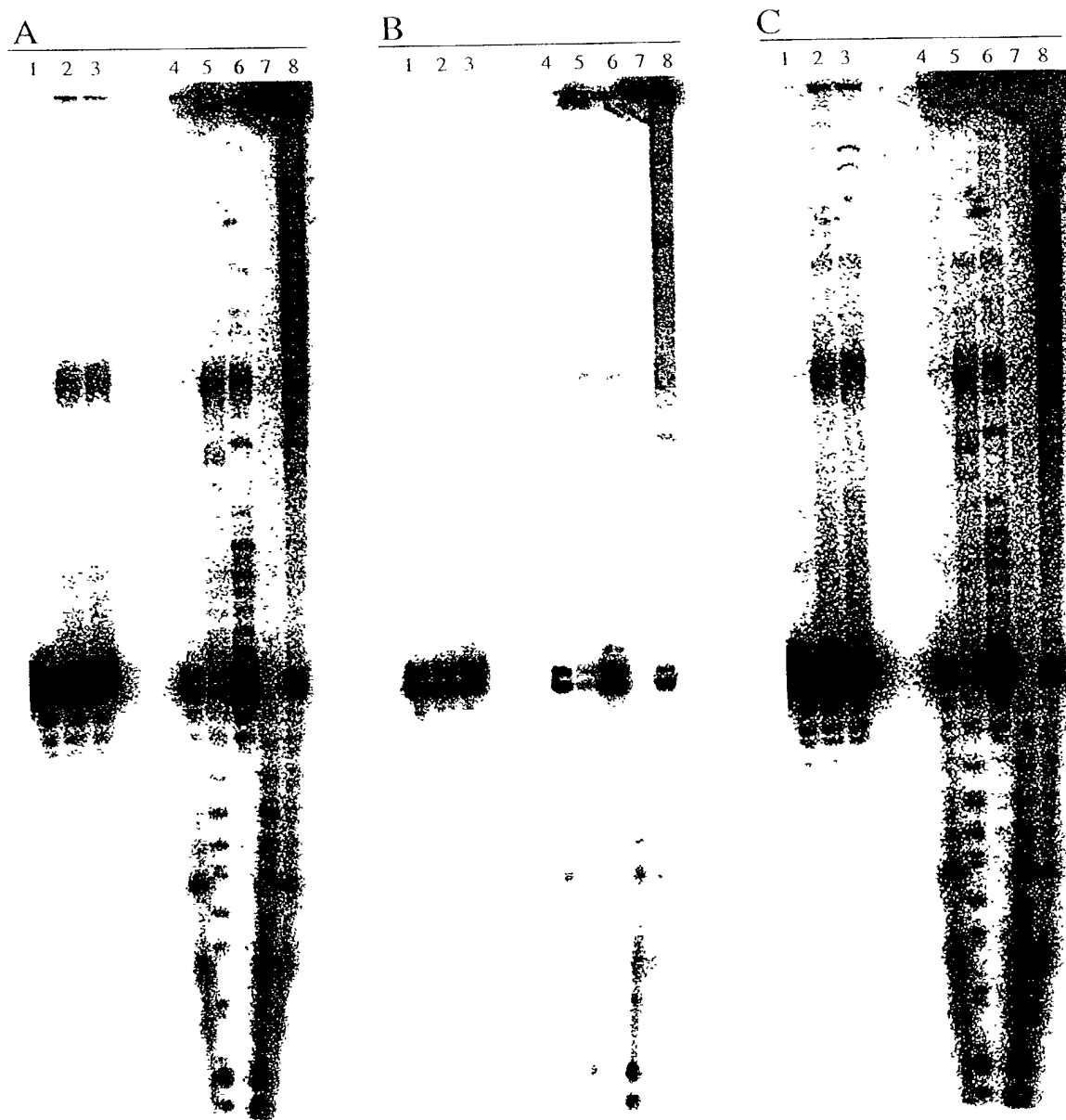
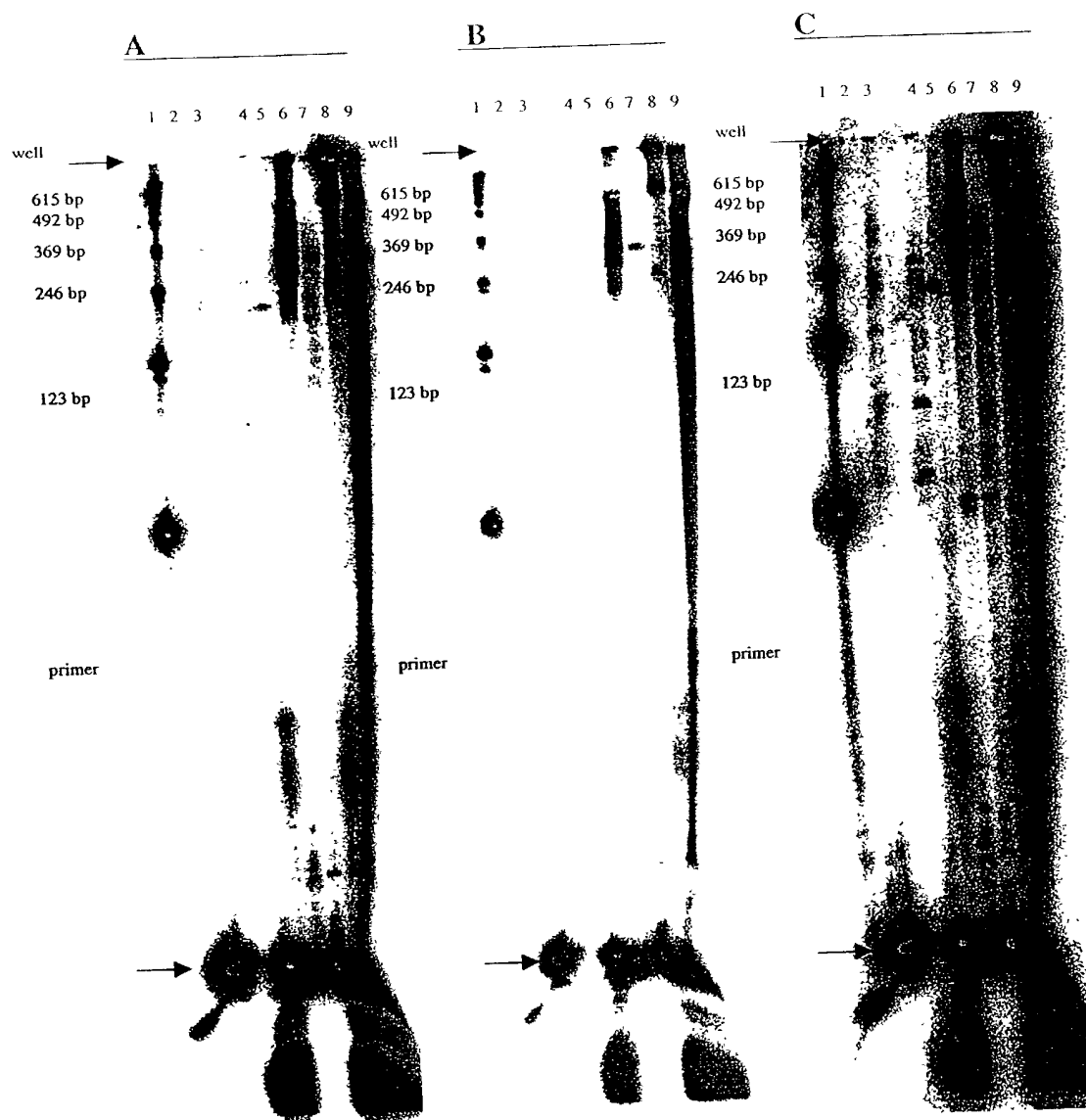


FIG. 6

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| | | Klenow | | | | | | | | Taq | |
|--------------|---|-----------------|----|------------|----------------|----|------------|------------------|----|-----------------|---------------|
| Enzyme | - | + | + | + | + | + | + | + | + | + | + |
| Primer (TOP) | + | + | + | + | + | + | + | + | + | + | + |
| Template | - | <u>BOT - 3T</u> | | | <u>BOT - T</u> | | | <u>BOT - Sau</u> | | <u>BOT - 3T</u> | |
| Nucleotide | - | dG | dA | γ A | dG | dA | γ A | dG | dA | γ A | dA γ A |

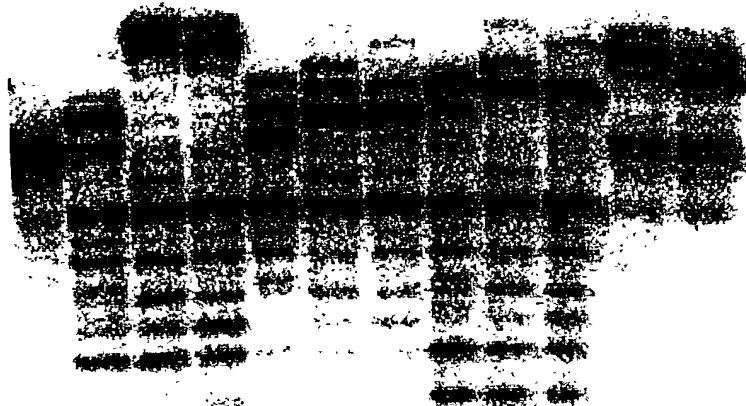


FIG. 8

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Pfu Primer Extension Assays

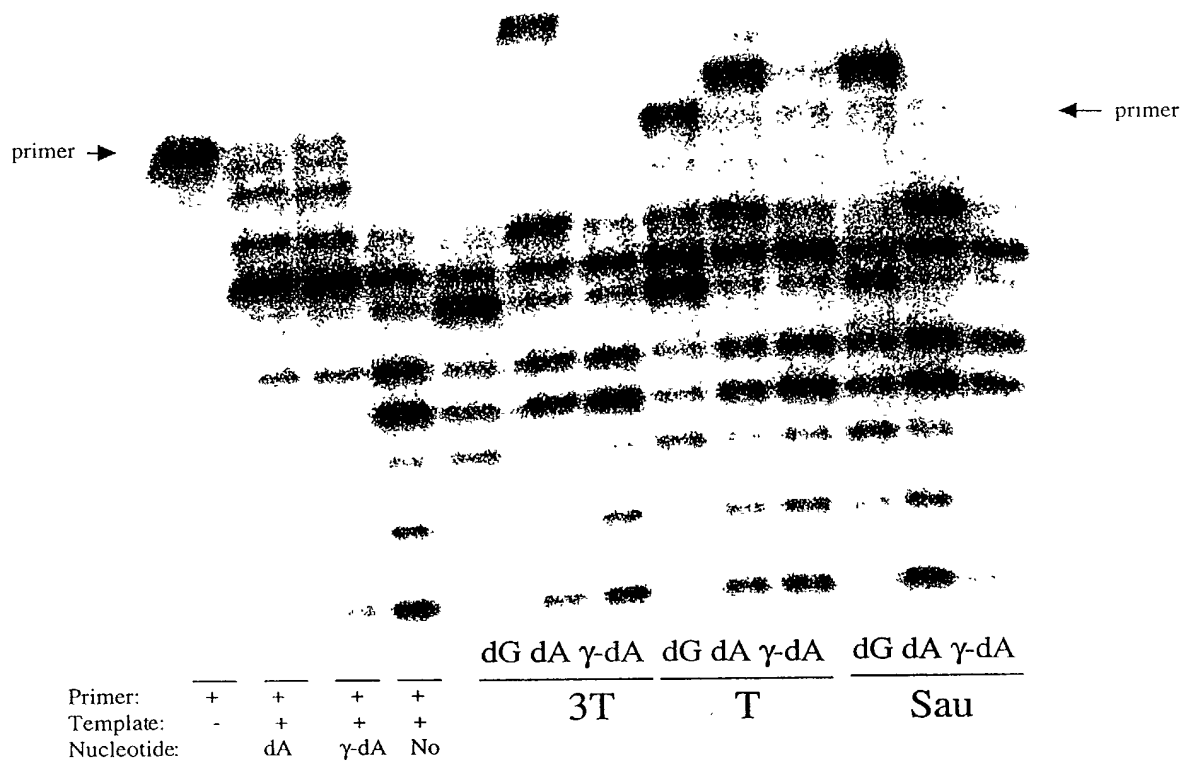


FIG. 9

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- Primer Strand:
Top 5' GGT ACT AAG CGG CCG CAT G 3'
- Template Strands:
3T 3' CCA TGA TTC GCC GGC GTA CTT TC 5'
Sau 3' CCA TGA TTC GCC GGC GTA CCT AG 5'

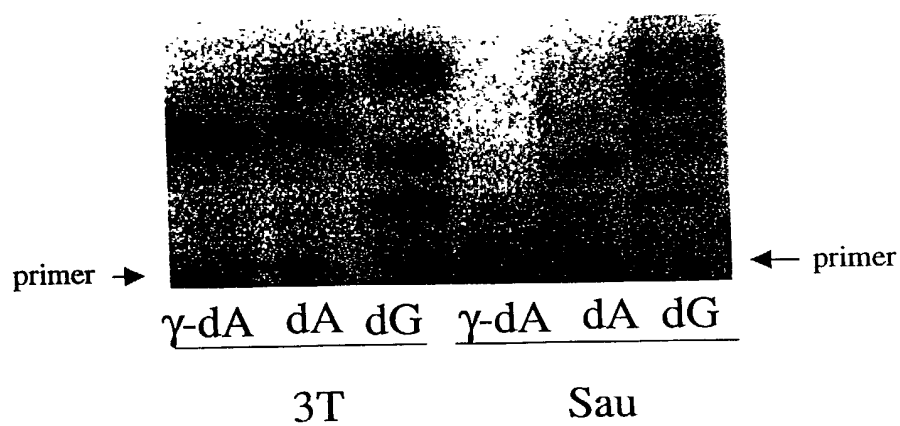


FIG. 10

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Primer Strand:

Top 5' GGT ACT AAG CGG CCG CAT G 3'

Template Strands:

BOT-3T 3' CCA TGA TTC GCC GGC GTA CTT TC 5'
 BOT-Sau 3' CCA TGA TTC GCC GGC GTA CCT AG 5'

| Enzyme: | None | T7 | T7 | Seq | Seq | T7 | | | | Sequenase | | | | Taq | | | |
|-------------|------|----|-------------|-----|-------------|----|----|---------|-------------|-----------|--------|---------------|-----|---------------|--------|-------------|-------------|
| Primer: | + | + | + | + | + | + | + | + | + | + | + | + | + | + | + | + | |
| Template: | - | + | - | + | BOT-3T | | | | Sau | | BOT-3T | | Sau | | BOT-3T | | |
| Nucleotide: | - | dA | γ dA | dA | γ dA | dG | dA | (spill) | γ dA | dG | dA | γ dAdG | dA | γ dAdG | dA | γ dA | γ dA |

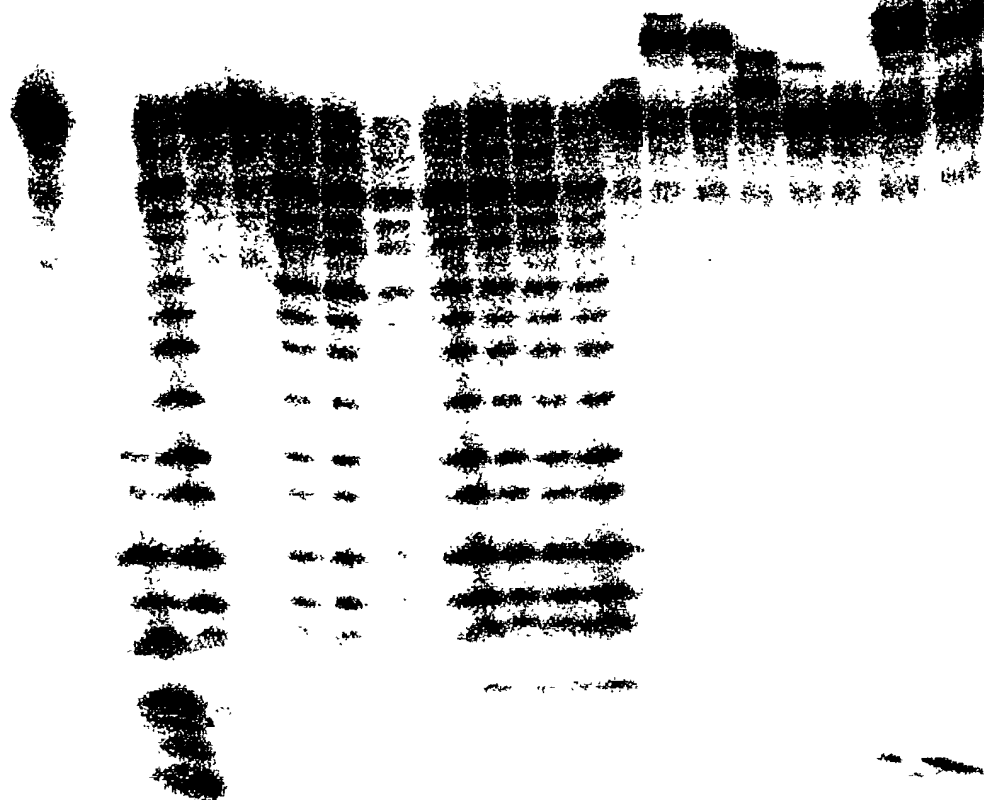
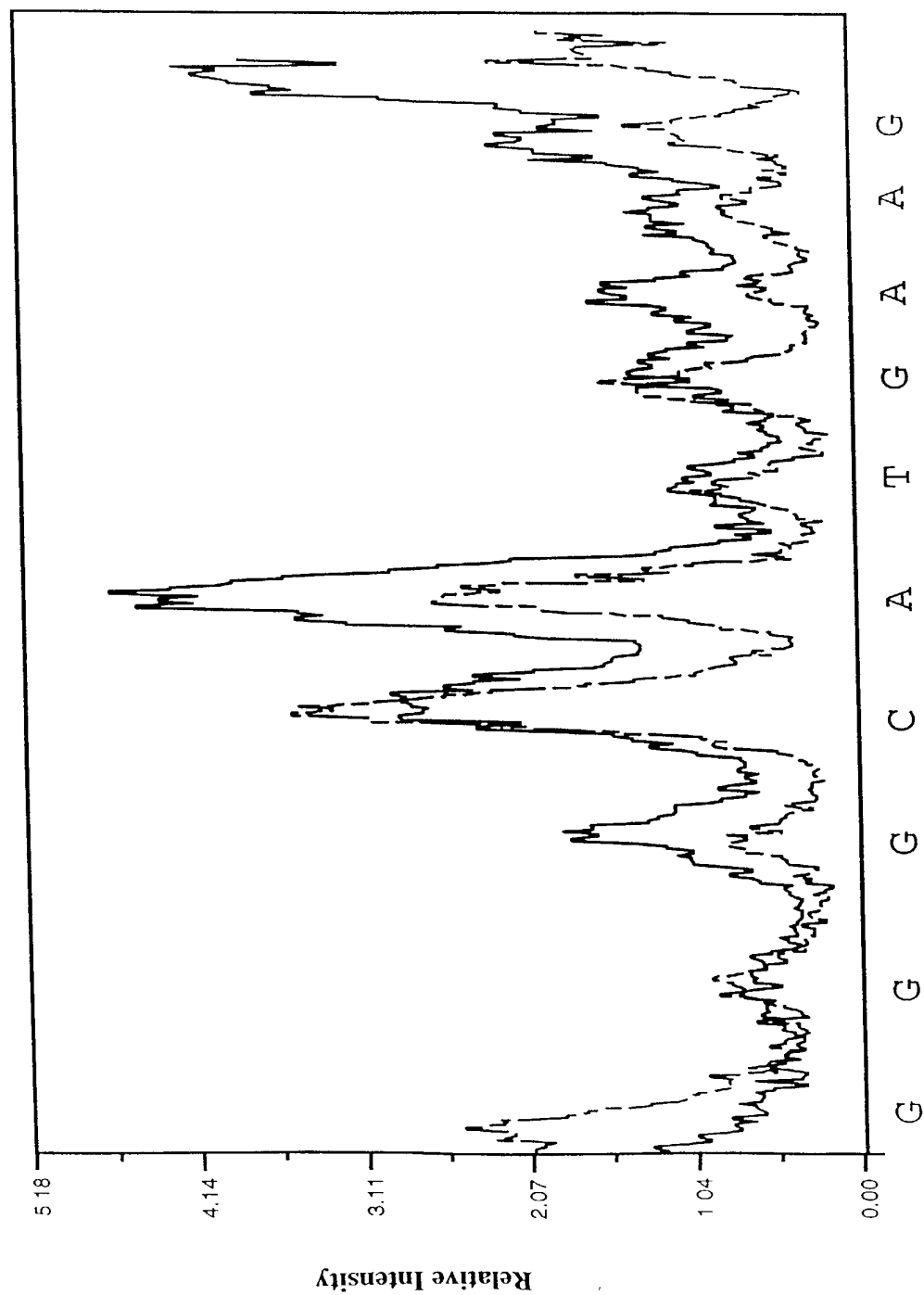


FIG. 11



Signal Intensity and Reaction Kinetics Provide Information Concerning Base Identity.

FIG. 12